ERROR DETECTED   SUGGESTED CORRECTION   SERIAL NUMBER: _09/787,995_

ATTN: NEW RULES CASES: PLEASE DISREGARD ENGLISH "ALPHA" HEADERS, WHICH WERE INSERTED BY PTO SOFTWARE

1____Wrapped Nucleics
     Wrapped Aminos

The number/text at the end of each line "wrapped" down to the next line. This may occur if your file was retrieved in a word processor after creating it. Please adjust your right margin to .3; this will prevent "wrapping."

2____Invalid Line Length

The rules require that a line not exceed 72 characters in length. This includes white spaces.

3____Misaligned Amino
     Numbering

The numbering under each 5th amino acid is misaligned. Do not use tab codes between numbers; use space characters, instead.

4____Non-ASCII

The submitted file was not saved in ASCII(DOS) text, as required by the Sequence Rules. Please ensure your subsequent submission is saved in ASCII text.

5____Variable Length

Sequence(s)____contain n's or Xaa's representing more than one residue. Per Sequence Rules, each n or Xaa can only represent a single residue. Please present the maximum number of each residue having variable length and indicate in the <220>-<223> section that some may be missing.

6____PatentIn 2.0
     "bug"

A "bug" in PatentIn version 2.0 has caused the <220>-<223> section to be missing from amino acid sequences(s) _____. Normally, PatentIn would automatically generate this section from the previously coded nucleic acid sequence. Please manually copy the relevant <220>-<223> section to the subsequent amino acid sequence. This applies to the mandatory <220>-<223> sections for Artificial or Unknown sequences.

7____Skipped Sequences
     (OLD RULES)

Sequence(s) _____ missing. If intentional, please insert the following lines for each skipped sequence:
(2) INFORMATION FOR SEQ ID NO:X: (insert SEQ ID NO where "X" is shown)
(i)   SEQUENCE CHARACTERISTICS: (Do not insert any subheadings under this heading)
(xi) SEQUENCE DESCRIPTION:SEQ ID NO:X: (insert SEQ ID NO where "X" is shown)
This sequence is intentionally skipped

Please also adjust the "(ii) NUMBER OF SEQUENCES:" response to include the skipped sequences.

8____Skipped Sequences
     (NEW RULES)

Sequence(s) _____ missing. If intentional, please insert the following lines for each skipped sequence.
<210> sequence id number
<400> sequence id number
000

9____Use of n's or Xaa's
     (NEW RULES)

Use of n's and/or Xaa's have been detected in the Sequence Listing.
Per 1.823 of Sequence Rules, use of <220>-<223> is MANDATORY if n's or Xaa's are present.
In <220> to <223> section, please explain location of n or Xaa, and which residue n or Xaa represents.

10____Invalid <213>
      Response

Per 1.823 of Sequence Rules, the only valid <213> responses are: Unknown, Artificial Sequence, or scientific name (Genus/species). <220>-<223> section is required when <213> response is Unknown or is Artificial Sequence

11____Use of <220>

Sequence(s) _____ missing the <220> "Feature" and associated numeric identifiers and responses.
Use of <220> to <223> is MANDATORY if <213> "Organism" response is "Artificial Sequence" or "Unknown." Please explain source of genetic material in <220> to <223> section.
(See "Federal Register," 06/01/1998, Vol. 63, No. 104, pp. 29631-32) (Sec. 1.823 of Sequence Rules)

12____PatentIn 2.0
      "bug"

Please do not use "Copy to Disk" function of PatentIn version 2.0. This causes a corrupted file, resulting in missing mandatory numeric identifiers and responses (as indicated on raw sequence listing). Instead, please use "File Manager" or any other manual means to copy file to floppy disk.

13____Misuse of n

n can only be used to represent a single nucleotide in a nucleic acid sequence. N is not used to represent any value not specifically a nucleotide.

AMC/MH – Biotechnology Systems Branch – 08/21/2001

PCT09

RAW SEQUENCE LISTING　　　　　　DATE: 01/22/2002
PATENT APPLICATION:　US/09/787,995　　TIME: 16:33:33

Input Set : A:\ST98032.ST25.txt
Output Set: N:\CRF3\01182002\I787995.raw

Does Not Comply
Corrected Diskette Need:

```
 3 <110> APPLICANT: Aventis Pharma S.A.
 5 <120> TITLE OF INVENTION: Use Of Specific Hybrid Promoters For Controlling Tissue
Expression
 7 <130> FILE REFERENCE: ST98032
 9 <140> CURRENT APPLICATION NUMBER: US/09/787,995
 9 <141> CURRENT FILING DATE: 2001-03-23
 9 <150> PRIOR APPLICATION NUMBER: US 60/123298
10 <151> PRIOR FILING DATE: 1999-04-04
12 <150> PRIOR APPLICATION NUMBER: FR/98/12000
13 <151> PRIOR FILING DATE: 1998-09-25
15 <160> NUMBER OF SEQ ID NOS: 10
17 <170> SOFTWARE: PatentIn version 3.0
19 <210> SEQ ID NO: 1
20 <211> LENGTH: 23
21 <212> TYPE: DNA
22 <213> ORGANISM: Artificial
24 <220> FEATURE:
25 <223> OTHER INFORMATION: Artificial Sequence
27 <220> FEATURE:
28 <221> NAME/KEY: nucleotide
29 <222> LOCATION: (1)..(23)
31 <400> SEQUENCE: 1
32 gatggtccct acttatgctg cta
35 <210> SEQ ID NO: 2
36 <211> LENGTH: 23
37 <212> TYPE: DNA
38 <213> ORGANISM: Artificial
40 <220> FEATURE:
41 <223> OTHER INFORMATION: articifial sequence
43 <220> FEATURE:
44 <221> NAME/KEY: nucleotide
45 <222> LOCATION: (1)..(23)
47 <400> SEQUENCE: 2
48 cttccatcat accaaactac ata
51 <210> SEQ ID NO: 3
52 <211> LENGTH: 32
53 <212> TYPE: DNA
54 <213> ORGANISM: Artificial
56 <220> FEATURE:
57 <223> OTHER INFORMATION: artificial sequence
59 <220> FEATURE:
60 <221> NAME/KEY: nucleotide
61 <222> LOCATION: (1)..(32)
63 <400> SEQUENCE: 3
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give source of genetic material – see item //

on Error Summary Sheet

23

misspelled　*see item //*

23

item //

```
                    RAW SEQUENCE LISTING          DATE: 01/22/2002
                    PATENT APPLICATION:  US/09/787,995   TIME: 16:33:33

                    Input Set : A:\ST98032.ST25.txt
                    Output Set: N:\CRF3\01182002\I787995.raw

       64 ctgctaaatt gctcgaggac aaattagaca aa                        32
       67 <210> SEQ ID NO: 4
       68 <211> LENGTH: 33
       69 <212> TYPE: DNA
C--> 70 <213> ORGANISM: Artificial
       72 <220> FEATURE:
       73 <223> OTHER INFORMATION: artificial sequence
       75 <220> FEATURE:
W--> 76 <221> NAME/KEY: oligonucleotide
       77 <222> LOCATION: (1)..(33)
       79 <400> SEQUENCE: 4
       80 ccctgacaaa gcttggctgg gctgctccac tgg                       33
       83 <210> SEQ ID NO: 5
       84 <211> LENGTH: 30
       85 <212> TYPE: DNA
C--> 86 <213> ORGANISM: Artificial
       88 <220> FEATURE:
       89 <223> OTHER INFORMATION: artificial sequence
       91 <220> FEATURE:
W--> 92 <221> NAME/KEY: oligonucleotide
       93 <222> LOCATION: (1)..(30)
       95 <400> SEQUENCE: 5
       96 atcgacgcgt gcccgttaca taacttacgg                           30
       99 <210> SEQ ID NO: 6
      100 <211> LENGTH: 38
      101 <212> TYPE: DNA
C--> 102 <213> ORGANISM: Artificial
      104 <220> FEATURE:
      105 <223> OTHER INFORMATION: artificial sequence
      107 <220> FEATURE:
W--> 108 <221> NAME/KEY: oligonucleotide
      109 <222> LOCATION: (1)..(38)
      111 <400> SEQUENCE: 6
      112 atcgacgcgt ccgctcgagc gtcaatgggg cggagttg                  38
      115 <210> SEQ ID NO: 7
      116 <211> LENGTH: 36
      117 <212> TYPE: DNA
C--> 118 <213> ORGANISM: Artificial
      120 <220> FEATURE:
      121 <223> OTHER INFORMATION: artificial sequence
      123 <220> FEATURE:
W--> 124 <221> NAME/KEY: oligonucleotide
      125 <222> LOCATION: (1)..(36)
      127 <400> SEQUENCE: 7
      128 ccaggctgca ctcgagacta gttcccacca actcga                    36
      131 <210> SEQ ID NO: 8
      132 <211> LENGTH: 36
      133 <212> TYPE: DNA
C--> 134 <213> ORGANISM: Artificial
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                    RAW SEQUENCE LISTING              DATE: 01/22/2002
                    PATENT APPLICATION:  US/09/787,995  TIME: 16:33:33

                    Input Set : A:\ST98032.ST25.txt
                    Output Set: N:\CRF3\01182002\I787995.raw

        136 <220> FEATURE:
        137 <223> OTHER INFORMATION: artificial sequence
        139 <220> FEATURE:
W--> 140 <221> NAME/KEY: oligonucleotide
        141 <222> LOCATION: (1)..(36)
        143 <400> SEQUENCE: 8
        144 tcgtttgaag cttggaagga gagtagcttc ggtgtc                    36
        147 <210> SEQ ID NO: 9
        148 <211> LENGTH: 30
        149 <212> TYPE: DNA
C--> 150 <213> ORGANISM: Artificial
        152 <220> FEATURE:
        153 <223> OTHER INFORMATION: artificial sequence
        155 <220> FEATURE:
W--> 156 <221> NAME/KEY: oligonucleotide
        157 <222> LOCATION: (1)..(30)
        159 <400> SEQUENCE: 9
        160 cccgttacat aacttacggt aaatggcccg                          30
        163 <210> SEQ ID NO: 10
        164 <211> LENGTH: 30
        165 <212> TYPE: DNA
C--> 166 <213> ORGANISM: Artificial
        168 <220> FEATURE:
        169 <223> OTHER INFORMATION: artificial sequence
        171 <220> FEATURE:
W--> 172 <221> NAME/KEY: oligonucleotide
        173 <222> LOCATION: (1)..(30)
        175 <400> SEQUENCE: 10
        176 gggacgcgct tctacaaggc gctggccgaa                          30
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VERIFICATION SUMMARY   DATE: 01/22/2002
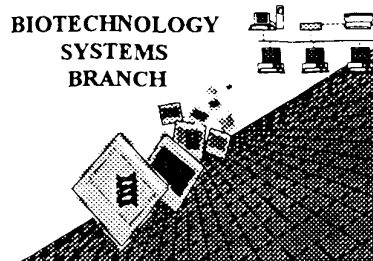PATENT APPLICATION: **US/09/787,995**   TIME: 16:33:34

Input Set : **A:\ST98032.ST25.txt**
Output Set: **N:\CRF3\01182002\I787995.raw**

L:9 M:270 C: Current Application Number differs, Replaced Current Application No
L:9 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:22 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:1
L:28 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:1
L:38 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:2
L:44 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:2
L:54 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:3
L:60 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:3
L:70 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:4
L:76 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:4
L:86 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:5
L:92 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:5
L:102 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:6
L:108 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:6
L:118 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:7
L:124 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:7
L:134 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:8
L:140 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:8
L:150 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:9
L:156 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:9
L:166 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:10
L:172 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:10

PK

RAW SEQUENCE LISTING
ERROR REPORT

The Biotechnology Systems Branch of the Scientific and Technical Information
Center (STIC) detected errors when processing the following computer readable
form:

Application Serial Number: _09/787,995_
Source: _PCT/09_
Date Processed by STIC: _1/22/2002_

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.
PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:
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   APPLICANT, WITH A NOTICE TO COMPLY or,
2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A
   NOTICE TO COMPLY
FOR CRF SUBMISSION QUESTIONS, PLEASE CONTACT MARK SPENCER, 703-308-4212.

FOR SEQUENCE RULES INTERPRETATION, PLEASE CONTACT ROBERT WAX, 703-308-4216.
PATENTIN 2.1 e-mail help: patin21help@uspto.gov or phone 703-306-4119 (R. Wax)
PATENTIN 3.0 e-mail help: patin3help@uspto.gov or phone 703-306-4119 (R. Wax)

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VERSION 3.1 PROGRAM**, ACCESSIBLE THROUGH THE U.S. PATENT AND
TRADEMARK OFFICE WEBSITE: SEE BELOW FOR ADDRESS:
      http://www.uspto.gov/web/offices/pac/checker

Applicants submitting genetic sequence information electronically on diskette or CD-Rom should be aware that there is
a possibility that the disk/CD-Rom may have been affected by treatment given to all incoming mail.
Please consider using alternate methods of submission for the disk/CD-Rom or replacement disk/CD-Rom, including:
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   User Manual** - ePAVE)
2. U.S. Patent and Trademark Office, Box Sequence, P.O. Box 2327, Arlington, VA 22202
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   U.S. Patent and Trademark Office, Technology Center 1600, Reception Area, 7th Floor, Examiner Name,
   Sequence Information, Crystal Mall One, 1911 South Clark Street, Arlington, VA 22202
            Or
   U.S. Patent and Trademark Office, Box Sequence, Customer Window, Lobby, Room 1B03, Crystal Plaza Two,
   2011 South Clark Place, Arlington, VA 22202
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